



ENTERED

PCT09

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/936,921B

DATE: 02/14/2003  
TIME: 13:24:51

Input Set : A:\110530.txt  
Output Set: N:\CRF4\02142003\I936921B.raw

3 <110> APPLICANT: RAOULT, Didier  
4 LA SCOLA, Bernard  
5 BIRG, Marie-Laure  
6 FENOLLAR, Florence  
8 <120> TITLE OF INVENTION: DIAGNOSIS OF WHIPPLE'S DISEASE  
10 <130> FILE REFERENCE: 110530  
12 <140> CURRENT APPLICATION NUMBER: 09/936,921B  
13 <141> CURRENT FILING DATE: 2001-09-24  
15 <150> PRIOR APPLICATION NUMBER: FR 99/03989  
16 <151> PRIOR FILING DATE: 1999-03-26  
18 <150> PRIOR APPLICATION NUMBER: FR 99/06679  
19 <151> PRIOR FILING DATE: 1999-05-21  
21 <150> PRIOR APPLICATION NUMBER: PCT/FR00/00754  
22 <151> PRIOR FILING DATE: 2000-03-24  
24 <160> NUMBER OF SEQ ID NOS: 5  
26 <170> SOFTWARE: PatentIn version 3.1  
28 <210> SEQ ID NO: 1  
29 <211> LENGTH: 19  
30 <212> TYPE: DNA  
31 <213> ORGANISM: Artificial Sequence  
33 <220> FEATURE:  
34 <223> OTHER INFORMATION: primer  
36 <220> FEATURE:  
37 <221> NAME/KEY: misc\_feature  
38 <222> LOCATION: (2)..(2)  
39 <223> OTHER INFORMATION: n = I  
42 <220> FEATURE:  
43 <221> NAME/KEY: misc\_feature  
44 <222> LOCATION: (8)..(8)  
45 <223> OTHER INFORMATION: n = I  
48 <220> FEATURE:  
49 <221> NAME/KEY: misc\_feature  
50 <222> LOCATION: (9)..(9)  
51 <223> OTHER INFORMATION: n = I  
54 <220> FEATURE:  
55 <221> NAME/KEY: misc\_feature  
56 <222> LOCATION: (11)..(11)  
57 <223> OTHER INFORMATION: n = I  
60 <220> FEATURE:  
61 <221> NAME/KEY: misc\_feature  
62 <222> LOCATION: (14)..(14)  
63 <223> OTHER INFORMATION: n = I  
66 <400> SEQUENCE: 1

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W--> 67 tnatgggnnc naanatgca 19  
 70 <210> SEQ ID NO: 2  
 71 <211> LENGTH: 20  
 72 <212> TYPE: DNA  
 73 <213> ORGANISM: Artificial Sequence  
 75 <220> FEATURE:  
 76 <223> OTHER INFORMATION: primer  
 78 <220> FEATURE:  
 79 <221> NAME/KEY: misc\_feature  
 80 <222> LOCATION: (6)..(6)  
 81 <223> OTHER INFORMATION: n = I  
 84 <220> FEATURE:  
 85 <221> NAME/KEY: misc\_feature  
 86 <222> LOCATION: (15)..(15)  
 87 <223> OTHER INFORMATION: n = I  
 90 <220> FEATURE:  
 91 <221> NAME/KEY: misc\_feature  
 92 <222> LOCATION: (18)..(18)  
 93 <223> OTHER INFORMATION: n = I  
 96 <400> SEQUENCE: 2  
 W--> 97 gcccancatt ccatntcncc 20  
 100 <210> SEQ ID NO: 3  
 101 <211> LENGTH: 612  
 102 <212> TYPE: DNA  
 103 <213> ORGANISM: Tropheyryma whippelii  
 105 <220> FEATURE:  
 106 <221> NAME/KEY: misc\_feature  
 107 <222> LOCATION: (9)..(9)  
 108 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
 111 <220> FEATURE:  
 112 <221> NAME/KEY: misc\_feature  
 113 <222> LOCATION: (16)..(16)  
 114 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
 117 <220> FEATURE:  
 118 <221> NAME/KEY: misc\_feature  
 119 <222> LOCATION: (18)..(18)  
 120 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
 123 <220> FEATURE:  
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 125 <222> LOCATION: (22)..(22)  
 126 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
 129 <220> FEATURE:  
 130 <221> NAME/KEY: misc\_feature  
 131 <222> LOCATION: (29)..(29)  
 132 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
 135 <220> FEATURE:  
 136 <221> NAME/KEY: misc\_feature  
 137 <222> LOCATION: (38)..(38)  
 138 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other

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141 <220> FEATURE:  
142 <221> NAME/KEY: misc\_feature  
143 <222> LOCATION: (68)..(68)  
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147 <220> FEATURE:  
148 <221> NAME/KEY: misc\_feature  
149 <222> LOCATION: (117)..(118)  
150 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
153 <220> FEATURE:  
154 <221> NAME/KEY: misc\_feature  
155 <222> LOCATION: (195)..(195)  
156 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
159 <220> FEATURE:  
160 <221> NAME/KEY: misc\_feature  
161 <222> LOCATION: (379)..(379)  
162 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
165 <220> FEATURE:  
166 <221> NAME/KEY: misc\_feature  
167 <222> LOCATION: (436)..(436)  
168 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown or other  
171 <400> SEQUENCE: 3  
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174 aattgccnag gatccggca ttgtggggca tggttccgct gatgttgta ctgtcannca 120  
176 ggatgacggg aaacatcgcg actaccatgt tggttaaattt gttcggttcaa atcaggggcaa 180  
178 ctgttacaac cagcnagatgg tggccgatc cggagatcgt gtataaaaag gtacagttct 240  
180 tgcacatggt ccatgtactg acaaagggtga gcttagtctt ggtagaaatc ttctgggtgc 300  
182 tttcatgccc tgggagggtc ataacttga ggtatgcata attatcagcc agaatttgg 360  
184 caaggacgac acccttcnt caatccacat agaagaacat gaggttagca cccgggatac 420  
186 gaagctggc agtganagaa ataacgcgag accttccgaa tgtaagcatg gattacataa 480  
188 aggacttgaa cgaacggggtt attatccgaa ttggcgctga ggttggccct gggacattt 540  
190 tggttggtaa ggtgacccca aagggcgaag accgaactca acgcggaaaga acgtttgctg 600  
192 agggctatct tt 612  
195 <210> SEQ ID NO: 4  
196 <211> LENGTH: 18  
197 <212> TYPE: DNA  
198 <213> ORGANISM: Artificial Sequence  
200 <220> FEATURE:  
201 <223> OTHER INFORMATION: oligonucleotide  
203 <400> SEQUENCE: 4 18  
204 gcattgtggg ggatgttt  
207 <210> SEQ ID NO: 5  
208 <211> LENGTH: 18  
209 <212> TYPE: DNA  
210 <213> ORGANISM: Artificial Sequence  
212 <220> FEATURE:  
213 <223> OTHER INFORMATION: oligonucleotide  
215 <400> SEQUENCE: 5  
216 ttgggggtcac cttaccaa 18

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/936,921B

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 2,8,9,11,14  
Seq#:2; N Pos. 6,15,18  
Seq#:3; N Pos. 9,16,18,22,29,38,68,117,118,195,379,436

**VERIFICATION SUMMARY**

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L:67 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

L:97 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0

L:172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0

M:341 Repeated in SeqNo=3